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(54) Title: HUMAN INTERLEUKIN-11 RECEPTOR (57) Abstract Polynucleotides encoding the human IL-11 receptor and fragments thereof are disclosed. IL-11 receptor proteins, methods for their production, inhibitors of binding of human IL-11 and its receptor and methods for their identification are also disclosed.		

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HUMAN INTERLEUKIN-11 RECEPTOR

Field of the Invention

5 The present invention relates to the human interleukin-11 receptor, fragments thereof and recombinant polynucleotides and cells useful for expressing such proteins.

Background of the Invention

10 A variety of regulatory molecules, known as cytokines, have been identified including interleukin-11 (IL-11). The various protein forms of IL-11 and DNA encoding various forms of IL-11 activity are described in Bennett, *et al.*, USPN 5,215,895 (June 1, 1993); McCoy, *et al.*, USPN 5,270,181 (December 14, 1993); and McCoy, *et al.*, USPN 5,292,646 (March 8, 1994),
15 all incorporated herein by reference. Thus, the term "IL-11" includes proteins having the biological activity described in these patents, whether produced by recombinant genetic engineering techniques; purified from cell sources producing the factor naturally or upon induction with other factors; or synthesized by chemical techniques; or a combination of the foregoing.

20 IL-11 is a pleiotropic cytokine that has been implicated in production of several biological activities including: induction of multipotential hematopoietic progenitor cell proliferation (Musashi *et al.* (1991) *Blood*, 78, 1448-1451); enhancement of megakaryocyte and platelet formation (Burstein *et al.* (1992) *J. Cell. Physiol.*, 153, 305-312); stimulation of acute phase protein synthesis

(Baumann et al. (1991) J. Biol. Chem., 266, 20424-20427); inhibition of adipocyte lipoprotein lipase activity (Kawashima et al. (1991) FEBS Lett., 283, 199-202); and effects on neurotransmitter phenotype (Fann et al. (1994) Proc. Natl. Acad. Sci. USA, 91, 43-47).

5 IL-11 may be used in a pharmaceutical preparation or formulation to treat immune deficiencies, specifically deficiencies in hematopoietic progenitor cells, or disorders relating thereto. Treatment of the other disorders or stimulation of the immune systems of cells thereof may also employ IL-11. IL-11 may also be employed in methods for treating cancer and other disease.

10 Such pathological states may result from disease, exposure to radiation or drugs, and include, for example, leukopenia, bacterial and viral infections, anemia, B cell or T cell deficiencies such as immune cell or hematopoietic cell deficiency following a bone marrow transplantation. IL-11 may also be used to potentiate the immune response to a variety of vaccines creating longer

15 lasting and more effective immunity. Therapeutic treatment of cancer and other diseases with IL-11 may avoid undesirable side effects caused by treatment with presently available drugs.

Like most cytokines, IL-11 exhibits certain biological activities by interacting with an IL-11 receptor (IL-11R) on the surface of target cells. It

20 would be desirable to identify and clone the sequence for the human receptor so that IL-11R proteins can be produced for various reasons, including production of therapeutics and screening for inhibitors of IL-11 binding to the receptor and receptor signalling.

Summary of the Invention

In accordance with the present invention, polynucleotides encoding the human interleukin-11 receptor are disclosed. In certain embodiments, the invention provides an isolated polynucleotide comprising a nucleotide sequence
5 selected from the group consisting of:

(a) the nucleotide sequence of SEQ ID NO:1 from nucleotide 803 to nucleotide 1999;

(b) a nucleotide sequence varying from the sequence of the nucleotide sequence specified in (a) as a result of degeneracy of the genetic
10 code; and

(c) an allelic variant of the nucleotide sequence specified in (a).

Preferably, the nucleotide sequence encodes a protein having a biological activity of the human IL-11 receptor. The nucleotide sequence may be operably linked to an expression control sequence. In preferred embodiments, the
15 polynucleotide comprises the nucleotide sequence of SEQ ID NO:1 from nucleotide 803 to nucleotide 1999 or a fragment thereof; the nucleotide sequence of SEQ ID NO:1 from nucleotide 803 to nucleotide 1828 or a fragment thereof; the nucleotide sequence of SEQ ID NO:1 from nucleotide 1904 to nucleotide 1999 or a fragment thereof; the nucleotide sequence of SEQ
20 ID NO:1 from nucleotide 734 to nucleotide 1999 or a fragment thereof; the nucleotide sequence of SEQ ID NO:1 from nucleotide 1067 to nucleotide 1828 or a fragment thereof; or the nucleotide sequence of SEQ ID NO:1 from nucleotide 1067 to nucleotide 1999 or a fragment thereof. In other embodiments, the polynucleotide comprises a nucleotide sequence capable of

hybridizing to the nucleotides sequence of SEQ ID NO:1 under highly stringent conditions.

The invention also provides isolated polynucleotides comprising a nucleotide sequence encoding a peptide or protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO:2;
- (b) the amino acid sequence of SEQ ID NO:2 from amino acids 24 to 422;
- (c) the amino acid sequence of SEQ ID NO:2 from amino acids 24 to 365;
- (d) the amino acid sequence of SEQ ID NO:2 from amino acids 391 to 422;
- (e) the amino acid sequence of SEQ ID NO:2 from amino acids 112 to 422;
- (f) the amino acid sequence of SEQ ID NO:2 from amino acids 112 to 365; and
- (g) fragments of (a)-(f) having a biological activity of the human IL-11 receptor.

Host cells, preferably mammalian cells, transformed with the polynucleotides are also provided.

In other embodiments, the invention provides a process for producing a human IL-11R protein. The process comprises:

- (a) growing a culture of the host cell of the present invention in a suitable culture medium; and

- (b) purifying the human IL-11R protein from the culture.

Proteins produced according to these methods are also provided.

The present invention also provides for an isolated human IL-11R protein comprising an amino acid sequence selected from the group consisting
5 of:

- (a) the amino acid sequence of SEQ ID NO:2;
- (b) the amino acid sequence of SEQ ID NO:2 from amino acids 24
to 422;
- (c) the amino acid sequence of SEQ ID NO:2 from amino acids 24
10 to 365;
- (d) the amino acid sequence of SEQ ID NO:2 from amino acids 391
to 422;
- (e) the amino acid sequence of SEQ ID NO:2 from amino acids 112
to 422;
- 15 (f) the amino acid sequence of SEQ ID NO:2 from amino acids 112
to 365; and
- (g) fragments of (a)-(f) having a biological activity of the human IL-
11 receptor. Preferably the protein comprises the amino acid sequence of SEQ
ID NO:2; the sequence from amino acid 24 to 422 of SEQ ID NO:2; the
20 sequence from amino acid 24 to 365 of SEQ ID NO:2; or the sequence from
amino acid 391 to 422 of SEQ ID NO:2. Pharmaceutical compositions
comprising a protein of the present invention and a pharmaceutically acceptable
carrier are also provided.

The present invention further provides for compositions comprising an antibody which specifically reacts with a protein of the present invention.

Methods of identifying an inhibitor of IL-11 binding to the human IL-11 receptor are also provided. These methods comprise:

- 5 (a) combining a human IL-11R protein or a fragment thereof with IL-11 or a fragment thereof, said combination forming a first binding mixture;
- (b) measuring the amount of binding between the protein and the IL-11 or fragment in the first binding mixture;
- (c) combining a compound with the protein and the IL-11 or
10 fragment to form a second binding mixture;
- (d) measuring the amount of binding in the second binding mixture;
and
- (e) comparing the amount of binding in the first binding mixture with
the amount of binding in the second binding mixture;
- 15 wherein the compound is capable of inhibiting IL-11 binding to the human IL-11 receptor when a decrease in the amount of binding of the second binding mixture occurs. Optionally, the first and/or second binding mixture may further comprise gp130 or a fragment thereof capable of binding to the protein of claim 11 or the IL-11 or fragment used therein. Inhibitors of IL-11R
20 identified by these methods and pharmaceutical compositions containing them are also provided.

Methods of inhibiting binding of IL-11 to the human IL-11 receptor in a mammalian subject are also disclosed which comprise administering a therapeutically effective amount of a composition containing a human IL-11R

protein, an IL-11R inhibitor or an antibody to a human IL-11R protein. Methods of treating or preventing loss of bone mass in a mammalian subject using these compositions are also provided.

5 Brief Description of the Figures

Figure 1 depicts a schematic representation of the structures of the human IL-11 receptor and gp130.

Figure 2 presents data demonstrating the biological activity of a soluble form of recombinant human IL-11R protein.

10

Detailed Description of Preferred Embodiments

The inventors of the present application have for the first time identified and provided a polynucleotide encoding the human IL-11 receptor (human IL-11R).

15 SEQ ID NO:1 provides the nucleotide sequence of a cDNA encoding the human IL-11R. SEQ ID NO:2 provides the amino acid sequence of the receptor, included a putative signal sequence from amino acids 1-23. The mature human IL-11R is believed to have the sequence of amino acids 24-422 of SEQ ID NO:2.

20 The mature receptor has at least three distinct domains: an extracellular domain (comprising approximately amino acids 24-365 of SEQ ID NO:2), a transmembrane domain (comprising approximately amino acids 366-390 of SEQ ID NO:2) and an intracellular domain (comprising approximately amino acids 391-422 of SEQ ID NO:2). The extracellular domain is further divided into an

immunoglobulin-like domain (comprising approximately amino acids 24-111 of SEQ ID NO:2) and a type-I-cytokine domain (comprising approximately amino acids 112-365 of SEQ ID NO:2).

5 Soluble forms of human IL-11R protein can also be produced. Such soluble forms include without limitation proteins comprising amino acids 1-365 and 24-365 of SEQ ID NO:2. The soluble forms of the human IL-11R are further characterized by being soluble in aqueous solution, preferably at room temperature. Human IL-11R proteins comprising only the intracellular domain or a portion thereof may also be produced. Any forms of human IL-11R of
10 less than full length are encompassed within the present invention and are referred to herein collectively as "human IL-11R" or "human IL-11R proteins." Human IL-11R proteins of less than full length may be produced by expressing a corresponding fragment of the polynucleotide encoding the full-length human IL-11R protein (SEQ ID NO:1). These corresponding polynucleotide fragments
15 are also part of the present invention. Modified polynucleotides as described above may be made by standard molecular biology techniques, including construction of appropriate desired deletion mutants, site-directed mutagenesis methods or by the polymerase chain reaction using appropriate oligonucleotide primers.

20 Based upon similarity to the structure of the IL-6 receptor, it is predicted that IL-11R proteins containing only the type-I cytokine domain of the extracellular region of the full length receptor will be capable of binding IL-11 and inducing receptor signalling. As a result, IL-11R proteins comprising amino acids 112 to 365 of SEQ ID NO:2, IL-11R proteins comprising amino

acids 112 to 390 of SEQ ID NO:2, and IL-11R proteins comprising amino acids 112 to 422 of SEQ ID NO:2 are provided by the present invention. Polynucleotides encoding such proteins (such as for example a polynucleotide comprising the nucleotides sequence of SEQ ID NO:1 from nucleotide 1067 to
5 nucleotide 1828, a polynucleotide comprising the nucleotides sequence of SEQ ID NO:1 from nucleotide 1067 to nucleotide 1906, and a polynucleotide comprising the nucleotides sequence of SEQ ID NO:1 from nucleotide 1067 to nucleotide 1999, respectively) are also provided by the invention.

For the purposes of the present invention, a protein has "a biological
10 activity of the human IL-11 receptor" if it possess one or more of the following characteristics: (1) the ability to bind IL-11 or a fragment thereof (preferably a biologically active fragment thereof); (2) the ability to bind to cytosolic proteins or molecules involved in the signalling pathway invoked by IL-11 binding to human IL-11R; (3) the ability to produce a signal characteristic of
15 the binding of IL-11 to human IL-11R (where the protein in question either contains a portion able to bind IL-11 or where the protein in question would produce such signal if joined to another protein which is able to bind IL-11); (4) the ability to bind to gp130 or a fragment thereof (either in the presence or absence of IL-11); (5) the ability to induce tyrosine phosphorylation of gp130;
20 (6) the ability to induce tyrosine phosphorylation of JAK kinases; or (7) the ability to induce tyrosine phosphorylation of the STAT family of DNA binding proteins (Zhong et al. (1994) Science 264, 95-98). Preferably, the biological activity possessed by the protein is the ability to bind IL-11 or a fragment

hereof, more preferably with a K_D of about 0.1 to about 100 nM, most preferably with a K_D of about 1 to about 10 nM.

Human IL-11R or active fragments thereof (human IL-11R proteins) may be fused to carrier molecules such as immunoglobulins. For example, soluble forms of the human IL-11R may be fused through "linker" sequences to the Fc portion of an immunoglobulin. Other fusion proteins, such as those with GST, Lex-A or MBP, may also be used.

The invention also encompasses allelic variations of the nucleotide sequence as set forth in SEQ ID NO:1, that is, naturally-occurring alternative forms of the isolated polynucleotide of SEQ ID NO: 1 which also encode human IL-11R proteins, preferably those proteins having a biological activity of human IL-11R. Also included in the invention are isolated polynucleotides which hybridize to the nucleotide sequence set forth in SEQ ID NO:1 under highly stringent conditions (for example, 0.1xSSC at 65°C). Isolated polynucleotides which encode human IL-11R proteins but which differ from the nucleotide sequence set forth in SEQ ID NO:1 by virtue of the degeneracy of the genetic code are also encompassed by the present invention. Variations in the nucleotide sequence as set forth in SEQ ID NO:1 which are caused by point mutations or by induced modifications are also included in the invention.

The isolated polynucleotides of the invention may be operably linked to an expression control sequence such as the pMT2 or pED expression vectors disclosed in Kaufman et al., Nucleic Acids Res. 19, 4485-4490 (1991), in order to produce the human IL-11R protein recombinantly. Many suitable expression control sequences are known in the art. General methods of expressing

recombinant proteins are also known and are exemplified in R. Kaufman, Methods in Enzymology 185, 537-566 (1990). As defined herein "operably linked" means enzymatically or chemically ligated to form a covalent bond between the isolated polynucleotide of the invention and the expression control
5 sequence, in such a way that the human IL-11R protein is expressed by a host cell which has been transformed (transfected) with the ligated polynucleotide/expression control sequence.

A number of types of cells may act as suitable host cells for expression of the human IL-11R protein. Any cell type capable of expressing functional
10 human IL-11R protein may be used. Suitable mammalian host cells include, for example, monkey COS cells, Chinese Hamster Ovary (CHO) cells, human kidney 293 cells, human epidermal A431 cells, human Colo205 cells, 3T3 cells, CV-1 cells, other transformed primate cell lines, normal diploid cells, cell strains derived from in vitro culture of primary tissue, primary explants, HeLa
15 cells, mouse L cells, BHK, HL-60, U937, HaK, Rat2, BaF3, 32D, FDCP-1, PC12, M1x or C2C12 cells.

The human IL-11R protein may also be produced by operably linking the isolated polynucleotide of the invention to suitable control sequences in one or more insect expression vectors, and employing an insect expression system.
20 Materials and methods for baculovirus/insect cell expression systems are commercially available in kit form from, e.g., Invitrogen, San Diego, California, U.S.A. (the MaxBac® kit), and such methods are well known in the art, as described in Summers and Smith, Texas Agricultural Experiment Station Bulletin No. 1555 (1987), incorporated herein by reference. Soluble forms of

the human IL-11R protein may also be produced in insect cells using appropriate isolated polynucleotides as described above.

Alternatively, the human IL-11R protein may be produced in lower eukaryotes such as yeast or in prokaryotes such as bacteria. Suitable yeast strains include *Saccharomyces cerevisiae*, *Schizosaccharomyces pombe*, Kluyveromyces strains, *Candida*, or any yeast strain capable of expressing heterologous proteins. Suitable bacterial strains include *Escherichia coli*, *Bacillus subtilis*, *Salmonella typhimurium*, or any bacterial strain capable of expressing heterologous proteins.

Expression in bacteria may result in formation of inclusion bodies incorporating the recombinant protein. Thus, refolding of the recombinant protein may be required in order to produce active or more active material. Several methods for obtaining correctly folded heterologous proteins from bacterial inclusion bodies are known in the art. These methods generally involve solubilizing the protein from the inclusion bodies, then denaturing the protein completely using a chaotropic agent. When cysteine residues are present in the primary amino acid sequence of the protein, it is often necessary to accomplish the refolding in an environment which allows correct formation of disulfide bonds (a redox system). General methods of refolding are disclosed in Kohno, Meth. Enzym., 185:187-195 (1990). EP 0433225 and copending application USSN 08/163,877 describe other appropriate methods.

The human IL-11R protein of the invention may also be expressed as a product of transgenic animals, e.g., as a component of the milk of transgenic

cows, goats, pigs, or sheep which are characterized by somatic or germ cells containing a polynucleotide sequence encoding the human IL-11R protein.

The human IL-11R protein of the invention may be prepared by growing a culture transformed host cells under culture conditions necessary to express
5 the desired protein. The resulting expressed protein may then be purified from the culture medium or cell extracts. Soluble forms of the human IL-11R protein of the invention can be purified from conditioned media. Membrane-bound forms of human IL-11R protein of the invention can be purified by preparing a total membrane fraction from the expressing cell and extracting the
10 membranes with a non-ionic detergent such as Triton X-100.

The human IL-11R protein can be purified using methods known to those skilled in the art. For example, the human IL-11R protein of the invention can be concentrated using a commercially available protein concentration filter, for example, an Amicon or Millipore Pellicon ultrafiltration
15 unit. Following the concentration step, the concentrate can be applied to a purification matrix such as a gel filtration medium. Alternatively, an anion exchange resin can be employed, for example, a matrix or substrate having pendant diethylaminoethyl (DEAE) or polyethyleimine (PEI) groups. The matrices can be acrylamide, agarose, dextran, cellulose or other types
20 commonly employed in protein purification. Alternatively, a cation exchange step can be employed. Suitable cation exchangers include various insoluble matrices comprising sulfopropyl or carboxymethyl groups. Sulfopropyl groups are preferred (e.g., S-Sepharose® columns). The purification of the human IL-11R protein from culture supernatant may also include one or more column

steps over such affinity resins as concanavalin A-agarose, heparin-toyopearl® or Cibacrom blue 3GA Sepharose®; or by hydrophobic interaction chromatography using such resins as phenyl ether, butyl ether, or propyl ether; or by immunoaffinity chromatography. Finally, one or more reverse-phase high performance liquid chromatography (RP-HPLC) steps employing hydrophobic RP-HPLC media, e.g., silica gel having pendant methyl or other aliphatic groups, can be employed to further purify the human IL-11R protein. Affinity columns including IL-11 or fragments thereof or including antibodies to the IL-11R protein can also be used in purification in accordance with known methods. Some or all of the foregoing purification steps, in various combinations or with other known methods, can also be employed to provide a substantially purified isolated recombinant protein. Preferably, the isolated human IL-11R protein is purified so that it is substantially free of other mammalian proteins.

Human IL-11R proteins of the invention may also be used to screen for agents which are capable of binding to human IL-11R or interfere with the binding of IL-11 to the human IL-11R (either the extracellular or intracellular domains) and thus may act as inhibitors of normal binding and cytokine action (IL-11R inhibitors). Binding assays using a desired binding protein, immobilized or not, are well known in the art and may be used for this purpose using the human IL-11R protein of the invention. Purified cell based or protein based (cell free) screening assays may be used to identify such agents. For example, human IL-11R protein may be immobilized in purified form on a carrier and binding to purified human IL-11R protein may be measured in the presence and in the absence of potential inhibiting agents. A suitable binding

assay may alternatively employ a soluble form of human IL-11R of the invention.

In such a screening assay, a first binding mixture is formed by combining IL-11 or a fragment thereof and human IL-11R protein, and the amount of binding in the first binding mixture (B_0) is measured. A second binding mixture is also formed by combining IL-11 or a fragment thereof, human IL-11R protein, and the compound or agent to be screened, and the amount of binding in the second binding mixture (B) is measured. The amounts of binding in the first and second binding mixtures are compared, for example, by performing a calculation of the ratio B/B_0 . A compound or agent is considered to be capable of inhibiting binding if a decrease in binding in the second binding mixture as compared to the first binding mixture is observed. Optionally, gp130 can be added to one or both of the binding mixtures. The formulation and optimization of binding mixtures is within the level of skill in the art, such binding mixtures may also contain buffers and salts necessary to enhance or to optimize binding, and additional control assays may be included in the screening assay of the invention.

Compounds found to reduce the binding activity of human IL-11R protein to IL-11 or its fragment to any degree, preferably by at least about 10%, more preferably greater than about 50% or more, may thus be identified and then secondarily screened in other binding assays and in vivo assays. By these means compounds having inhibitory activity for IL-11R binding which may be suitable as therapeutic agents may be identified.

Human IL-11R proteins, and polynucleotides encoding them, may also be used as diagnostic agents for detecting the expression or presence of IL-11R, IL-11 or cells expressing IL-11R or IL-11. The proteins or polynucleotides may be employed for such purpose in standard procedures for diagnostics assays using these types of materials. Suitable methods are well known to those skilled in the art.

Human IL-11R acts as a mediator of the known biological activities of IL-11. As a result, isolated human IL-11R protein and IL-11R inhibitors may be useful in treatment or modulation of various medical conditions in which IL-11 is implicated or which are effected by the activity (or lack thereof) of IL-11 (collectively "IL-11-related conditions"). IL-11-related conditions include without limitation immune deficiencies, specifically deficiencies in hematopoietic progenitor cells, or disorders relating thereto, cancer and other disease. Such pathological states may result from disease, exposure to radiation or drugs, and include, for example, leukopenia, bacterial and viral infections, anemia, B cell or T cell deficiencies such as immune cell or hematopoietic cell deficiency following a bone marrow transplantation.

It is also believed that IL-11 and IL-11R may play a role in the regulation of bone maturation and repair (Girasole et al. (1994) J. Clin. Invest. 93, 1516-1524; Passeri et al. (1992) J. Bone Miner. Res., 7(S1), S110 Abst.; Passeri et al. (1993) J. Bone Miner. Res., 8(S1), S162 Abst.). As a result, human IL-11R protein and IL-11R inhibitors may be useful in treatment of bone loss (including that associated with osteoporosis, post-menopausal osteoporosis.

senile osteoporosis, idiopathic osteoporosis, Pagets disease, multiple myeloma, and hypogonadal conditions).

Human IL-11R protein and IL-11R inhibitors, purified from cells or recombinantly produced, may be used as a pharmaceutical composition when
5 combined with a pharmaceutically acceptable carrier. Such a composition may contain, in addition to human IL-11R or ligand and carrier, diluents, fillers, salts, buffers, stabilizers, solubilizers, and other materials well known in the art. The term "pharmaceutically acceptable" means a non-toxic material that does not interfere with the effectiveness of the biological activity of the active
10 ingredient(s). The characteristics of the carrier will depend on the route of administration.

The pharmaceutical composition of the invention may also contain cytokines, lymphokines, or other hematopoietic factors such as M-CSF, GM-CSF, IL-1, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-11, IL-12,
15 IL-15, G-CSF, stem cell factor, and erythropoietin. The pharmaceutical composition may contain thrombolytic or anti-thrombotic factors such as plasminogen activator and Factor VIII. The pharmaceutical composition may further contain other anti-inflammatory agents. Such additional factors and/or agents may be included in the pharmaceutical composition to produce a
20 synergistic effect with isolated human IL-11R protein or IL-11R inhibitor, or to minimize side effects caused by the isolated human IL-11R or IL-11R inhibitor. Conversely, isolated human IL-11R or IL-11R inhibitor may be included in formulations of the particular cytokine, lymphokine, other hematopoietic factor, thrombolytic or anti-thrombotic factor, or anti-

inflammatory agent to minimize side effects of the cytokine, lymphokine, other hematopoietic factor, thrombolytic or anti-thrombotic factor, or anti-inflammatory agent.

5 The pharmaceutical composition of the invention may be in the form of a liposome in which isolated human IL-11R protein or IL-11R inhibitor is combined, in addition to other pharmaceutically acceptable carriers, with amphipathic agents such as lipids which exist in aggregated form as micelles, insoluble monolayers, liquid crystals, or lamellar layers which in aqueous solution. Suitable lipids for liposomal formulation include, without limitation,
10 monoglycerides, diglycerides, sulfatides, lysolecithin, phospholipids, saponin, bile acids, and the like. Preparation of such liposomal formulations is within the level of skill in the art, as disclosed, for example, in U.S. Patent No. 4,235,871; U.S. Patent No. 4,501,728; U.S. Patent No. 4,837,028; and U.S. Patent No. 4,737,323, all of which are incorporated herein by reference.

15 As used herein, the term "therapeutically effective amount" means the total amount of each active component of the pharmaceutical composition or method that is sufficient to show a meaningful patient benefit, e.g., amelioration of symptoms of, healing of, or increase in rate of healing of such conditions. When applied to an individual active ingredient, administered alone, the term
20 refers to that ingredient alone. When applied to a combination, the term refers to combined amounts of the active ingredients that result in the therapeutic effect, whether administered in combination, serially or simultaneously.

In practicing the method of treatment or use of the present invention, a therapeutically effective amount of isolated human IL-11R protein or IL-11R

inhibitor is administered to a mammal. Isolated human IL-11R protein or IL-11R inhibitor may be administered in accordance with the method of the invention either alone or in combination with other therapies such as treatments employing cytokines, lymphokines or other hematopoietic factors. When co-
5 administered with one or more cytokines, lymphokines or other hematopoietic factors, human IL-11R protein or IL-11R inhibitor may be administered either simultaneously with the cytokine(s), lymphokine(s), other hematopoietic factor(s), thrombolytic or anti-thrombotic factors, or sequentially. If administered sequentially, the attending physician will decide on the appropriate
10 sequence of administering human IL-11R protein or IL-11R inhibitor in combination with cytokine(s), lymphokine(s), other hematopoietic factor(s), thrombolytic or anti-thrombotic factors.

Administration of human IL-11R protein or IL-11R inhibitor used in the pharmaceutical composition or to practice the method of the present invention
15 can be carried out in a variety of conventional ways, such as oral ingestion, inhalation, or cutaneous, subcutaneous, or intravenous injection. Intravenous administration to the patient is preferred.

When a therapeutically effective amount of human IL-11R protein or IL-11R inhibitor is administered orally, human IL-11R protein or IL-11R inhibitor
20 will be in the form of a tablet, capsule, powder, solution or elixir. When administered in tablet form, the pharmaceutical composition of the invention may additionally contain a solid carrier such as a gelatin or an adjuvant. The tablet, capsule, and powder contain from about 5 to 95% human IL-11R protein or IL-11R inhibitor, and preferably from about 25 to 90% human IL-11R

protein or IL-11R inhibitor. When administered in liquid form, a liquid carrier such as water, petroleum, oils of animal or plant origin such as peanut oil, mineral oil, soybean oil, or sesame oil, or synthetic oils may be added. The liquid form of the pharmaceutical composition may further contain physiological
5 saline solution, dextrose or other saccharide solution, or glycols such as ethylene glycol, propylene glycol or polyethylene glycol. When administered in liquid form, the pharmaceutical composition contains from about 0.5 to 90% by weight of human IL-11R protein or IL-11R inhibitor, and preferably from about 1 to 50% human IL-11R protein or IL-11R inhibitor.

10 When a therapeutically effective amount of human IL-11R protein or IL-11R inhibitor is administered by intravenous, cutaneous or subcutaneous injection, human IL-11R protein or IL-11R inhibitor will be in the form of a pyrogen-free, parenterally acceptable aqueous solution. The preparation of such parenterally acceptable protein solutions, having due regard to pH, isotonicity,
15 stability, and the like, is within the skill in the art. A preferred pharmaceutical composition for intravenous, cutaneous, or subcutaneous injection should contain, in addition to human IL-11R protein or IL-11R inhibitor an isotonic vehicle such as Sodium Chloride Injection, Ringer's Injection, Dextrose Injection, Dextrose and Sodium Chloride Injection, Lactated Ringer's Injection,
20 or other vehicle as known in the art. The pharmaceutical composition of the present invention may also contain stabilizers, preservatives, buffers, antioxidants, or other additive known to those of skill in the art.

The amount of human IL-11R protein or IL-11R inhibitor in the pharmaceutical composition of the present invention will depend upon the nature

and severity of the condition being treated, and on the nature of prior treatments which the patient has undergone. Ultimately, the attending physician will decide the amount of human IL-11R protein or IL-11R inhibitor with which to treat each individual patient. Initially, the attending physician will administer
5 low doses of human IL-11R protein or IL-11R inhibitor and observe the patient's response. Larger doses of human IL-11R protein or IL-11R inhibitor may be administered until the optimal therapeutic effect is obtained for the patient, and at that point the dosage is not generally increased further. It is contemplated that the various pharmaceutical compositions used to practice the
10 method of the present invention should contain about 0.1 μ g to about 100 mg of human IL-11R protein or IL-11R inhibitor per kg body weight.

The duration of intravenous therapy using the pharmaceutical composition of the present invention will vary, depending on the severity of the disease being treated and the condition and potential idiosyncratic response of
15 each individual patient. It is contemplated that the duration of each application of the human IL-11R protein or IL-11R inhibitor will be in the range of 12 to 24 hours of continuous intravenous administration. Ultimately the attending physician will decide on the appropriate duration of intravenous therapy using the pharmaceutical composition of the present invention.

20 Human IL-11R proteins of the invention may also be used to immunize animals to obtain polyclonal and monoclonal antibodies which specifically react with the human IL-11R protein and which may inhibit binding of IL-11 or fragments thereof to the receptor. Such antibodies may be obtained using the entire human IL-11R as an immunogen, or by using fragments of human IL-

11R, such as the soluble mature human IL-11R. Smaller fragments of the human IL-11R may also be used to immunize animals. The peptide immunogens additionally may contain a cysteine residue at the carboxyl terminus, and are conjugated to a hapten such as keyhole limpet hemocyanin (KLH). Additional peptide immunogens may be generated by replacing tyrosine residues with sulfated tyrosine residues. Methods for synthesizing such peptides are known in the art, for example, as in R.P. Merrifield, J. Amer. Chem. Soc. 85, 2149-2154 (1963); J.L. Krstenansky, et al., FEBS Lett. 211, 10 (1987).

Neutralizing or non-neutralizing antibodies (preferably monoclonal antibodies) binding to human IL-11R protein may also be useful therapeutics for certain tumors and also in the treatment of conditions described above. These neutralizing monoclonal antibodies may be capable of blocking IL-11 binding to the human IL-11R.

15

Example 1

Isolation of Human IL-11R cDNA

Generation of DNA Probes:

DNA probes derived from the murine Etl-2 sequence (SEQ ID NO:3) were obtained by PCR from murine placenta cDNA. The amino terminal probe corresponds to base pairs 418-570 and the carboxy terminal probe to base pairs 847- 1038 of the murine Etl-2 sequence. The DNA probes were gel purified and radiolabeled using $\alpha^{32}\text{P}$ -dATP and $\alpha^{32}\text{P}$ -dCTP.

cDNA Library Screening:

cDNA was generated from activated human PBMC using the Superscript Choice System and cloned into the EcoR1 site of ZAP II (Stratagene). The resulting phage were used to infect E. coli strain BB4. One million phage were
5 plated on 150 mm NZCYM plates at a density of 15000 pfu/plate. Plaques were transferred to duplicate Duralose nitrocellulose filters (Stratagene). Following alkali denaturation and heat fixation the filters were pre-hybridized in 5X SSC, 5X Denhardt's, 0.1% SDS, and 50 µg/ml yeast tRNA for 2 hours at 65°C. One set of filters was hybridized with the amino-terminal probe and
10 the other set with the carboxy-terminal probe (5×10^5 cpm/ml) for 48 hrs at 55°C in pre-hybridization buffer. The filters were washed with 4X SSC, 0.1% SDS once at 25°C and twice at 55°C. Plaques that hybridized to both probes were identified by autoradiography.

Of the one million plaques screened two plaques hybridized to both of
15 the probes. These plaques were picked and the phage eluted into SM media containing chloroform. The resulting phage were used to reinfect E. coli strain BB4 and plated on NZCYM plates at a density of 100-300 pfu/plate for a secondary screen.

Following the secondary screen plasmid DNA was isolated from the
20 ZAPII plaques by excision using helper phage (Stratagene). The DNA sequence of the inserts was determined on an Applied Biosystems DNA sequencer.

Clone phIL11R14-2 containing the polynucleotide having the sequence of SEQ ID NO:1 was deposited with ATCC at accession number _____ on December 22, 1994.

5

Example 2

Expression of Soluble Human IL-11R Protein and

Assay of Activity

A soluble form of human IL-11R protein was expressed in mammalian cells. The expressed recombinant protein was capable of transducing a signal
10 in BAF130-9 cells.

A portion of the full length human IL-11R sequence (nucleotides 734-1828 of SEQ ID NO:1 encoding amino acids 1-365 of SEQ ID NO:2) corresponding to a soluble form was cloned into the mammalian expression vector pED and used to transfect COSM6 cells. 40 hours after transfection
15 conditioned media was removed, concentrated 5 fold and used in proliferation assays with the murine cell line BAF130-9 (Hibi, M. et al. (1990) Cell 63, 1149-57), a derivative of the BAFB03 cell line expressing the human gp130 signal transducer. BAF130-9 cells do not proliferate in response to IL-11 or IL-6 alone, but do proliferate in response to a combination of IL-6 and soluble
20 IL-6R (Hibi et al., supra). BAF130-9 cells (1×10^4 cell in 0.1ml) were cultured in RPMI 1640 medium/10% FCS with increasing concentrations of recombinant human IL-11 in the absence or presence of 10 μ l of conditioned media from mock transfected cells or cells transfected with the soluble human IL-11R sequence. After forty hours the cells were pulse-labeled with ^3H -thymidine (0.5

5 Other human IL-11R proteins can be tested in this model to determine whether they exhibit a "biological activity" of human IL-11R as defined herein.

Other Systems for Determination Biological Activity of Human IL-11R

Protein

15 Assays for IL-11 Binding

Binding of IL-11 to the extracellular region of the human IL-11R protein

20 will specifically cause a rapid induction of phosphotyrosine on the receptor protein. Assays for ligand binding activity as measured by induction of phosphorylation are described below.

25

For example, a DNA construct is prepared in which the extracellular domain (truncated prior, preferably immediately prior, to the predicted transmembrane domain) is ligated in frame to a cDNA encoding the hinge C_H2 and C_H3 domains of a human immunoglobulin (Ig) γ 1. This construct is generated in an appropriate expression vector for COS cells, such as pED Δ C or pMT2. The
5 plasmid is transiently transfected into COS cells. The secreted IL-11R-Ig fusion protein is collected in the conditioned medium and purified by protein A chromatography.

The purified IL-11R-Ig fusion protein is used to demonstrate IL-11
10 binding in a number of applications. IL-11 can be coated onto the surface of an enzyme-linked immunosorbent assay (ELISA) plate, and then additional binding sites blocked with bovine serum albumin or casein using standard ELISA buffers. The IL-11R-Ig fusion protein is then bound to the solid-phase IL-11, and binding is detected with a secondary goat anti-human Ig conjugated
15 to horseradish peroxidase. The activity of specifically bound enzyme can be measured with a colorimetric substrate, such as tetramethyl benzidine and absorbance readings.

IL-11 may also be expressed on the surface of cells. for example by providing a transmembrane domain or glucosyl phosphatidyl inositol (GPI)
20 linkage. Cells expressing the membrane bound IL-11 can be identified using the IL-11R-Ig fusion protein. The soluble IL-11R-Ig fusion is bound to the surface of these cells and detected with goat anti-human Ig conjugated to a fluorochrome, such as fluorescein isothiocyanate and flow cytometry.

Interaction Trap

A yeast genetic selection method, the "interaction trap" [Gyuris et al, Cell 75:791-803, 1993], can be used to determine whether a human IL-11R protein has a biological activity of human IL-11R as defined herein. In this system, the expression of reporter genes from both LexAop-Leu2 and LexAop-LacZ relies on the interaction between the bait protein, for example in this case a species which interacts with human IL-11R, and the prey, for example in this case the human IL-11R protein. Thus, one can measure the strength of the interaction by the level of Leu2 or LacZ expression. The most simple method is to measure the activity of the LacZ encoded protein, β -galactosidase. This activity can be judged by the degree of blueness on the X-Gal containing medium or filter. For the quantitative measurement of β -galactosidase activity, standard assays can be found in "Methods in Yeast Genetics" Cold Spring Harbor, New York, 1990 (by Rose, M.D., Winston, F., and Hieter, P.).

In such methods, if one wishes to determine whether the human IL-11R protein interacts with a particular species (such as, for example, a cystolic protein which binds to the intracellular domain of the human IL-11R *in vivo*), that species can be used as the "bait" in the interaction trap with the human IL-11R protein to be tested serving as the "prey", or *vice versa*.

CAT Induction System

Transcription of acute phase plasma protein genes, such as the rat β -fibrinogen gene, is activated by IL-11 in a variety of cell lines. In one

exemplary system, COSM6 cells are cotransfected with plasmids encoding the human IL-11R protein (such as the full length human IL-11R or a soluble form thereof), the human gp130 signal transducer and a reporter gene containing the 350 base pair promoter region of the rat b-fibrinogen gene fused to a reporter gene, CAT (Baumann et al. (1991) J. Biol. Chem. 266, 20424-27). The cells are stimulated with increasing concentrations of recombinant human IL-11 and transcription of the reporter gene is monitored by assaying for the presence of CAT activity.

10 Phosphorylation of gp130

Activity may also be determined by examining the ability of IL-11 to induce tyrosine phosphorylation of gp130 in cells transfected with a sequence encoding the human IL-11R protein (such as the full length human IL-11R or a soluble form thereof) (Luttcken et al. (1994) Science 263, 89-92).

15

Phosphorylation of STATs

Activity may also be determined by examining the ability of IL-11 to induce tyrosine phosphorylation of STATs (signal transducers and activators of transcription, a family of DNA binding proteins) in cells transfected with a sequence encoding the human IL-11R protein (such as the full length human IL-11R or a soluble form thereof) (Zhong et al. (1994) Science 264, 95-98).

20

Phosphorylation of JAK Kinases

Activity may also be determined by examining the ability of IL-11 to induce tyrosine phosphorylation of JAK kinases in cells transfected with a sequence encoding the human IL-11R protein (such as the full length human IL-11R or a soluble form thereof) (Yin et al. (1993) J. Immunol. 151: 2555-61).

5

All patent and literature references cited herein are incorporated by reference as if fully set forth.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Tobin, James
- (ii) TITLE OF INVENTION: HUMAN INTERLEUKIN-11 RECEPTOR
- (iii) NUMBER OF SEQUENCES: 4
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Genetics Institute, Inc.
 - (B) STREET: 87 CambridgePark Drive
 - (C) CITY: Cambridge
 - (D) STATE: MA
 - (E) COUNTRY: USA
 - (F) ZIP: 02140

(v) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

- (A) APPLICATION NUMBER:
- (B) FILING DATE:
- (C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

- (A) NAME: Brown, Scott A.
- (B) REGISTRATION NUMBER: 32,724
- (C) REFERENCE/DOCKET NUMBER: GI5252

(ix) TELECOMMUNICATION INFORMATION:

- (A) TELEPHONE: (617) 498-8224
- (B) TELEFAX: (617) 876-5851

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 2456 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: double
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 734..1999

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

TGCCCCACCC	CCAGCCTCTG	GCAGCAGCCA	GGGCATCTGG	ATCTGCTTAA	CTACACAGCC	60
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CCAGGCCCCAG	CTGAGTGGCC	CAAGTAGCCA	GACCAAGGAG	CCAGGTTTCAG	GCGAGAAGCC	180
TGGCAGCCAG	GGCAGGGGTG	GGCCTCAGGG	TGGGAGTGCA	GGATGGGCTC	AGATCCATGA	240
TGACACCCCT	CCCCCAGGGT	GATAAGGTCT	GCCTAGGTTA	ATCAGAGGCA	GTGATAAGCC	300
CTGGACCAGG	TGGGGGTAAA	TACCAGAAAT	CCCAACAGCT	GGACTGGAGG	GGTTAATGGG	360
AGTGGCTGAG	CTGGTGCCAG	TGCTTGGTGC	CAGGGGTGGG	CGCCAAGGGC	AGTGGAGGGG	420
GAGTTGCTGG	CACAGTCTGT	TGCCTCCGGC	TTTTGTTCTG	GGCCCTAAGC	CCAGGACTGA	480
GATGGAGGGT	GTGAGGGGGT	GTGTGTGTCC	GTGTGTGTGT	GTGTGTGTGT	GTGCGCGCGC	540
ACGCACATGC	AAAGCACTGG	GTATACAGTG	GGAAAGGGGA	CCTCAGGTCA	GTTCCCGCAG	600
TGATTTCTAA	CAGCCTTACC	CCACTTGGTG	CATCAATTTT	TCTCCTAGGA	AGCCTCAGTT	660
TTGGAGAGGA	AGAGCCAGGC	TTTAGCCTCC	CATCTCAGGG	GTGCGGGATT	TTTGACTCTA	720

Sequence	Position	Residue
Met Ser Ser Ser Cys Ser Gly Leu Ser Arg Val Leu	1-10	
CTG GCC GTG GCT ACA GCC CTG GTG TCT GCC TCC TCC CCC TGC CCC CAG	11-20	
Val Ala Val Ala Thr Ala Leu Val Ser Ala Ser Ser Pro Cys Pro Gln	21-30	
GCC TGG GGC CCC CCA GGG GTC CAG TAT GGG CAG CCA GGC AGG TCC GTG	31-40	
Ala Trp Gly Pro Pro Gly Val Gln Tyr Gly Gln Pro Gly Arg Ser Val	41-50	
AAG CTG TGT TGT CCT GGA GTG ACT GCC GGG GAC CCA GTG TCC TGG TTT	51-60	
Lys Leu Cys Cys Pro Gly Val Thr Ala Gly Asp Pro Val Ser Trp Phe	61-70	
CGG GAT GGG GAG CCA AAG CTG CTC CAG GGA CCT GAC TCT GGG CTA GGG	71-80	
Arg Asp Gly Glu Pro Lys Leu Leu Gln Gly Pro Asp Ser Gly Leu Gly	81-90	
CAT GAA CTG GTC CTG GCC CAG GCA GAC AGC ACT GAT GAG GGC ACC TAC	91-100	
His Glu Leu Val Leu Ala Gln Ala Asp Ser Thr Asp Glu Gly Thr Tyr	101-110	
ATC TGC CAG ACC CTG GAT GGT GCA CTT GGG GGC ACA GTG ACC CTG CAG	111-120	
Ile Cys Gln Thr Leu Asp Gly Ala Leu Gly Gly Thr Val Thr Leu Gln	121-130	
CTG GGC TAC CCT CCA GCC CGC CCT GTT GTC TCC TGC CAA GCA GCC GAC	131-140	
Leu Gly Tyr Pro Pro Ala Arg Pro Val Val Ser Cys Gln Ala Ala Asp	141-150	
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CCC ACC CGC TAC CTC ACC TCC TAC AGG AAG ACA GTC CTA GGA GCT	171-180	
Pro Thr Arg Tyr Leu Thr Ser Tyr Arg Lys Lys Thr Val Leu Gly Ala	181-190	

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 Asp Pro Leu Gly Ala Ala Arg Cys Val Val His Gly Ala Glu Phe Trp 185
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 AGC CAG TAC CGG ATT AAT GTG ACT GAG GTG AAC CCA CTG GGT GCC AGC 1345
 Ser Gln Tyr Arg Ile Asn Val Thr Glu Val Asn Pro Leu Gly Ala Ser 200
 190
 ACA CGC CTG CTG GAT GTG AGC TTG CAG AGC ATC TTG CGC CCT GAC CCA 1393
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 CCC CAG GGC CTG CGG GTA GAG TCA GTA CCA GGT TAC CCC CGA CGC CTG 1441
 Pro Gln Gly Leu Arg Val Glu Ser Val Pro Gly Tyr Pro Arg Arg Leu 235
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 CGA GCC AGC TGG ACA TAC CCT GCC TCC TGG CCG TGC CAG CCC CAC TTC 1489
 Arg Ala Ser Trp Thr Tyr Pro Ala Ser Trp Pro Cys Gln Pro His Phe 250
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 CTG CTC AAG TTC CGT TTG CAG TAC CGT CCG GCG CAG CAT CCA GCC TGG 1537
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 255
 TCC ACG GTG GAG CCA GCT GGA CTG GAG GAG GTG ATC ACA GAT GCT GTG 1585
 Ser Thr Val Glu Pro Ala Gly Leu Glu Glu Val Ile Thr Asp Ala Val 280
 270
 GCT GGG CTG CCC CAT GCT GTA CGA GTC AGT GCC CGG GAC TTT CTA GAT 1633
 Ala Gly Leu Pro His Ala Val Arg Val Ser Ala Arg Asp Phe Leu Asp 300
 285
 GCT GGC ACC TGG AGC ACC TGG AGC CCG GAG GCC TGG GGA ACT CCG AGC 1681
 Ala Gly Thr Trp Ser Thr Trp Ser Pro Glu Ala Trp Gly Thr Pro Ser 315
 305
 ACT GGG ACC ATA CCA AAG GAG ATA CCA GCA TGG GGC CAG CTA CAC ACG 1729
 Thr Gly Thr Ile Pro Lys Glu Ile Pro Ala Trp Gly Gln Leu His Thr

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Gln Pro Glu Val Glu Pro Gln Val Asp Ser Pro Ala Pro Pro Arg Pro			
335	340	345	
TCC CTC CAA CCA CAC CCT CGG CTA CTT GAT CAC AGG GAC TCT GTG GAG			1825
Ser Leu Gln Pro His Pro Arg Leu Leu Asp His Arg Asp Ser Val Glu			
350	355	360	
CAG GTA GCT GTG CTG GCG TCT TTG GGA ATC CTT TCT TTC CTG GGA CTG			1873
Gln Val Ala Val Leu Ala Ser Leu Gly Ile Leu Ser Phe Leu Gly Leu			
365	370	375	380
GTG GCT GGG GCC CTG GCA CTG GGG CTC TGG CTG AGG CTG AGA CGG GGT			1921
Val Ala Gly Ala Leu Ala Leu Gly Leu Trp Leu Arg Leu Arg Arg Gly			
385	390	395	
GGG AAG GAT GGA TCC CCA AAG CCT GGG TTC TTG GCC TCA GTG ATT CCA			1969
Gly Lys Asp Gly Ser Pro Lys Pro Gly Phe Leu Ala Ser Val Ile Pro			
400	405	410	
GTG GAC AGG CGT CCA GGA GCT CCA AAC CTG TAGAGGACCC AGGAGGGCTT			2019
Val Asp Arg Arg Pro Gly Ala Pro Asn Leu			
415	420		
CGGCAGATTC CACCTATAAT TCTGTCTTGC TGGTGTGGAT GGATGGACAG ATAGAAACCA			2079
GGCAGGACAG TAGATCCCTA TGGTTGGATC TCAGCTGGAA GTTCTGTTTG GAGCCCATTT			2139
CTGTGAGACC CTGTATTTC AATTTCAGC TGAAGGGTGC TTGTACCTCT GATTTCACCC			2199
CAGAGTTGGA GTTCTGCTCA AGGAACGTGT GTAATGTGTA CATCTGTGTC CATGTGTGAC			2259
CATGTGTCTG TGAGGCAGGG AACATGTATT CTCTGCATGC ATGTATGTAG GTGCCCTGGG			2319
AGTGTGTGTG GGTCCCTTGGC TCTTGGCCTT TCCCCTTGCA GGGTTGTGC AGGTGTGAAT			2379
AAAGAGAATA AGGAAGTTCT TGGAGATTAT ACTCAGAAAA AAAAAAAAAA AGTCGACGCG			2439
GCCGCGAATT CCTGCAG			2456

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 422 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

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 35      40      45
Pro Gly Val Thr Ala Gly Asp Pro Val Ser Trp Phe Arg Asp Gly Glu
 50      55      60
Pro Lys Leu Leu Gln Gly Pro Asp Ser Gly Leu Gly His Glu Leu Val
 65      70      75      80
Leu Ala Gln Ala Asp Ser Thr Asp Glu Gly Thr Tyr Ile Cys Gln Thr
 85      90      95
Leu Asp Gly Ala Leu Gly Gly Thr Val Thr Leu Gln Leu Gly Tyr Pro
100      105      110
Pro Ala Arg Pro Val Val Ser Cys Gln Ala Ala Asp Tyr Glu Asn Phe
115      120      125
Ser Cys Thr Trp Ser Pro Ser Gln Ile Ser Gly Leu Pro Thr Arg Tyr
130      135      140
Leu Thr Ser Tyr Arg Lys Lys Thr Val Leu Gly Ala Asp Ser Gln Arg
145      150      155      160

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Arg Ser Pro Ser Thr Gly Pro Trp Pro Cys Pro Gln Asp Pro Leu Gly
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 Ile Asn Val Thr Glu Val Asn Pro Leu Gly Ala Ser Thr Arg Leu Leu
 195 200 205
 Asp Val Ser Leu Gln Ser Ile Leu Arg Pro Asp Pro Pro Gln Gly Leu
 210 215 220
 Arg Val Glu Ser Val Pro Gly Tyr Pro Arg Arg Leu Arg Ala Ser Trp
 225 230 235 240
 Thr Tyr Pro Ala Ser Trp Pro Cys Gln Pro His Phe Leu Leu Lys Phe
 245 250 255
 Arg Leu Gln Tyr Arg Pro Ala Gln His Pro Ala Trp Ser Thr Val Glu
 260 265 270
 Pro Ala Gly Leu Glu Glu Val Ile Thr Asp Ala Val Ala Gly Leu Pro
 275 280 285
 His Ala Val Arg Val Ser Ala Arg Asp Phe Leu Asp Ala Gly Thr Trp
 290 295 300
 Ser Thr Trp Ser Pro Glu Ala Trp Gly Thr Pro Ser Thr Gly Thr Ile
 305 310 315 320
 Pro Lys Glu Ile Pro Ala Trp Gly Gln Leu His Thr Gln Pro Glu Val
 325 330 335
 Glu Pro Gln Val Asp Ser Pro Ala Pro Pro Arg Pro Ser Leu Gln Pro
 340 345 350
 His Pro Arg Leu Leu Asp His Arg Asp Ser Val Glu Gln Val Ala Val
 355 360 365
 Leu Ala Ser Leu Gly Ile Leu Ser Phe Leu Gly Leu Val Ala Gly Ala
 370 375 380

Leu Ala Leu Gly Leu Trp Leu Arg Leu Arg Arg Gly Gly Lys Asp Gly
 385 390 395 400
 Ser Pro Lys Pro Gly Phe Leu Ala Ser Val Ile Pro Val Asp Arg Arg
 405 410 415
 Pro Gly Ala Pro Asn Leu
 420

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1714 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 34..1359

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

TTCTTAGCCT GATAGGAGGA AGTCTGGAG GCC ATG GCA CTC AGT CAC TGT GAT 54
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 Tyr Gln Asp Glu Gln Gln Leu Leu Arg Ala Asp Gln Gly Pro Gly Gly 20
 CGT GCT ACA GCC CTG GTG TCT TCC TCC TCC CCC TGC CCC CAA GCT TGG 150
 Arg Ala Thr Ala Leu Val Ser Ser Pro Cys Pro Gln Ala Trp 30
 25 35

GGT CCT CCA GGG GTC CAG TAT GGA CAA CCT GGC AGG CCC GTG ATG CTG 198
 Gly Pro Pro Gly Val Gln Tyr Gly Gln Pro Gly Arg Pro Val Met Leu 55
 40 45 50 55
 TGC TGC CCC GGA GTG AGT GCT GGG ACT CCA GTG TCC TGG TTT CGG GAT 246
 Cys Cys Pro Gly Val Ser Ala Gly Thr Pro Val Ser Trp Phe Arg Asp 70
 60 65 70
 GGA GAT TCA AGG CTG CTC CAG GGA CCT GAC TCT GGG TTA GGA CAC AGA 294
 Gly Asn Ser Arg Leu Leu Gln Gly Pro Asp Ser Gly Leu Gly His Arg 85
 75 80 85
 CTG GTC TTG GCC CAG GTG GAC AGC CCT GAT GAA GGC ACT TAT GTC TGC 342
 Leu Val Leu Ala Gln Val Asp Ser Pro Asp Glu Gly Thr Tyr Val Cys 100
 90 95 100
 CAG ACC CTG GAT GGT GTA TCA GGG GGC ATG GTG ACC CTG AAG CTG GGC 390
 Gln Thr Leu Asp Gly Val Ser Gly Gly Met Val Thr Leu Lys Leu Gly 115
 105 110 115
 TTT CCC CCA GCA CGT CCT GAA GTC TCC TGC CAA GCG GTA GAC TAT GAA 438
 Phe Pro Pro Ala Arg Pro Glu Val Ser Cys Gln Ala Val Asp Tyr Glu 135
 120 125 130 135
 AAC TTC TCC TGT ACT TGG AGT CCA GGC CAG GTC AGC GGT TTG CCC ACC 486
 Asn Phe Ser Cys Thr Trp Ser Pro Gly Gln Val Ser Gly Leu Pro Thr 150
 140 145 150
 CGC TAC CTT ACT TCC TAC AGG AAG AAG AAG AAG AAG AAG AAG AAG AAG 534
 Arg Tyr Leu Thr Ser Tyr Arg Lys Lys Thr Leu Pro Gly Ala Glu Ser 165
 155 160 165
 CAG AGG GAA AGT CCA TCC ACC GGG CCT TGG CCG TGT CCA CAG GAC CCT 582
 Gln Arg Glu Ser Pro Ser Thr Gly Pro Trp Pro Cys Pro Gln Asp Pro 180
 170 175 180
 CTG GAG GCC TCC CGA TGT GTG GTC CAT GGG GCA GAG TTC TGG AGT GAG 630
 Leu Glu Ala Ser Arg Cys Val Val His Gly Ala Glu Phe Trp Ser Glu 195
 185 190 195

TAC CGG ATC AAT GTG ACC GAG GTG AAC CCA CTG GGT GCC AGC ACG TGC 678
 Tyr Arg Ile Asn Val Thr Glu Val Asn Pro Leu Gly Ala Ser Thr Cys 215
 200 205
 CTA CTG GAT GTG AGA TTA CAG AGC ATC TTG CGT CCT GAT CCA CCC CAA 726
 Leu Leu Asp Val Arg Leu Gln Ser Ile Leu Arg Pro Asp Pro Pro Gln 230
 220 225
 GGA CTG CGG GTG GAA TCC GTA CCT GGT TAC CCG AGA CGC CTG CAT GCC 774
 Gly Leu Arg Val Glu Ser Val Pro Gly Tyr Pro Arg Arg Leu His Ala 245
 235 240
 AGC TGG ACA TAC CCT GCC TCC TGG CGT CGC CAA CCC CAC TTT CTG CTC 822
 Ser Trp Thr Tyr Pro Ala Ser Trp Arg Arg Gln Pro His Phe Leu Leu 260
 250 255
 AAG TTC CGG TTG CAA TAC CGA CCA GCA CAG CAT CCA GCC TGG TCC ACG 870
 Lys Phe Arg Leu Gln Tyr Arg Pro Ala Gln His Pro Ala Trp Ser Thr 275
 265 270
 GTG GAG CCC ATT GGC TTG GAG GAA GTG ATA ACA GAT GCT GTG GCT GGG 918
 Val Glu Pro Ile Gly Leu Glu Glu Val Ile Thr Asp Ala Val Ala Gly 295
 280 285
 CTG CCA CAC GCG GTA CGA GTC AGT GCC AGG GAC TTT CTG GAT GCT GGC 966
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 300 305
 ACC TGG AGC GCC TGG AGC CCA GAG GCC TGG GGT ACT CCT AGC ACT GGT 1014
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 315 320
 CCC CTG CAG GAT GAG ATA CCT GAT TGG AGC CAG GGA CAT GGA CAG CAG 1062
 Pro Leu Gln Asp Glu Ile Pro Asp Trp Ser Gln Gly His Gly Gln Gln 340
 330 335
 CTA GAG GCA GTA GCT CAG GAG GAC AGC CCG GCT CCT GCA AGG CCT 1110
 Leu Glu Ala Val Val Ala Gln Glu Asp Ser Pro Ala Pro Ala Arg Pro 355
 345 350
 TCC TTG CAG CCG GAC CCA AGG CCA CTT GAT CAC AGG GAC CCC TTG GAG 1158
 Ser Leu Gln Pro Asp Pro Arg Pro Leu Asp His Arg Asp Pro Leu Glu

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360          365          370          375          1206
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Gln Val Ala Val Leu Ala Ser Leu Gly Ile Phe Ser Cys Leu Gly Leu
380          385          390
GCT GTT GGA GCT CTG GCA CTG GGG CTC TGG CTG AGG CTG AGA CGG AGT
Ala Val Gly Ala Leu Ala Leu Gly Leu Trp Leu Arg Leu Arg Ser
395          400          405          1254
GGG AAG GAT GGA CCG CAA AAA CCT GGG CTC TTG GCA CCC ATG ATC CCG
Gly Lys Asp Gly Pro Gln Lys Pro Gly Leu Ala Pro Met Ile Pro
410          415          420          1302
GTG GAA AAG CTT CCA GGA ATT CCA AAC CTG CAG AGG ACC CCA GAG AAC
Val Glu Lys Leu Pro Gly Ile Pro Asn Leu Gln Arg Thr Pro Glu Asn
425          430          435          1350
TTC AGC TGATTTTCATC TGTAACCCGG TCAGACTTGG GGTGGTTAAA AGGACAGGCA
Phe Ser
440          1406
GAAAGAGGCG GGGCAGTGGA TCCCTGTGGA TGGAGGTCTC AGCTGAAAGT CTGAGCTCTT
1466
TTCTTTTGACA CCTATACTCC AAACCTTGCTG CCGGCTGAAG GCTGTCTGGA CTTCCGATGT
1526
CCTGAGGTGG AAGTCCACCT GAGGAATGTG TACAGAAAGTC TGTGTTCCCTG TGATCGTGTG
1586
TGTATGTGAG ACAGGGAGCA AAAGTTCTCT GCATGTGTGT ACAGATGATT GGAGAGTGTG
1646
TGCGGTCTTG GGCTTGGCCC TTCTGGGAAG TGTGAAGAGT TGAATAAAAA GAGACGGAAG
1706
TTTTTTGGA
1714

```

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 441 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Met Ala Leu Ser His Cys Asp Tyr Gln Asp Glu Gln Gln Leu Leu Arg
 1 5 10 15
 Ala Asp Gln Gly Pro Gly Gly Arg Ala Thr Ala Leu Val Ser Ser Ser
 20 25 30
 Ser Pro Cys Pro Gln Ala Trp Gly Pro Pro Gly Val Gln Tyr Gly Gln
 35 40 45
 Pro Gly Arg Pro Val Met Leu Cys Cys Pro Gly Val Ser Ala Gly Thr
 50 55 60
 Pro Val Ser Trp Phe Arg Asp Gly Asp Ser Arg Leu Leu Gln Gly Pro
 65 70 75 80
 Asp Ser Gly Leu Gly His Arg Leu Val Leu Ala Gln Val Asp Ser Pro
 85 90 95
 Asp Glu Gly Thr Tyr Val Cys Gln Thr Leu Asp Gly Val Ser Gly Gly
 100 105 110
 Met Val Thr Leu Lys Leu Gly Phe Pro Pro Ala Arg Pro Glu Val Ser
 115 120 125
 Cys Gln Ala Val Asp Tyr Glu Asn Phe Ser Cys Thr Trp Ser Pro Gly
 130 135 140
 Gln Val Ser Gly Leu Pro Thr Arg Tyr Leu Thr Ser Tyr Arg Lys Lys
 145 150 155 160
 Thr Leu Pro Gly Ala Glu Ser Gln Arg Glu Ser Pro Ser Thr Gly Pro
 165 170 175
 Trp Pro Cys Pro Gln Asp Pro Leu Glu Ala Ser Arg Cys Val Val His
 180 185 190
 Gly Ala Glu Phe Trp Ser Glu Tyr Arg Ile Asn Val Thr Glu Val Asn
 195 200 205

Pro Leu Gly Ala Ser Thr Cys Leu Leu Asp Val Arg Leu Gln Ser Ile
 210 215 220
 Leu Arg Pro Asp Pro Pro Gln Gly Leu Arg Val Glu Ser Val Pro Gly
 225 230 235 240
 Tyr Pro Arg Arg Leu His Ala Ser Trp Thr Tyr Pro Ala Ser Trp Arg
 245 250 255
 Arg Gln Pro His Phe Leu Leu Lys Phe Arg Leu Gln Tyr Arg Pro Ala
 260 265 270
 Gln His Pro Ala Trp Ser Thr Val Glu Pro Ile Gly Leu Glu Glu Val
 275 280 285
 Ile Thr Asp Ala Val Ala Gly Leu Pro His Ala Val Arg Val Ser Ala
 290 295 300
 Arg Asp Phe Leu Asp Ala Gly Thr Trp Ser Ala Trp Ser Pro Glu Ala
 305 310 315 320
 Trp Gly Thr Pro Ser Thr Gly Pro Leu Gln Asp Glu Ile Pro Asp Trp
 325 330 335
 Ser Gln Gly His Gly Gln Gln Leu Glu Ala Val Ala Gln Glu Asp
 340 345 350
 Ser Pro Ala Pro Ala Arg Pro Ser Leu Gln Pro Asp Pro Arg Pro Leu
 355 360 365
 Asp His Arg Asp Pro Leu Glu Gln Val Ala Val Leu Ala Ser Leu Gly
 370 375 380
 Ile Phe Ser Cys Leu Gly Leu Ala Val Gly Ala Leu Ala Leu Gly Leu
 385 390 395 400
 Trp Leu Arg Leu Arg Arg Ser Gly Lys Asp Gly Pro Gln Lys Pro Gly
 405 410 415
 Leu Leu Ala Pro Met Ile Pro Val Glu Lys Leu Pro Gly Ile Pro Asn
 420 425 430

Leu Gln Arg Thr Pro Glu Asn Phe Ser
435 440

What is claimed is:

1. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:1 from nucleotide 803 to nucleotide 1999;
 - (b) a nucleotide sequence varying from the sequence of the nucleotide sequence specified in (a) as a result of degeneracy of the genetic code; and
 - (c) an allelic variant of the nucleotide sequence specified in (a).
2. The polynucleotide of claim 1 wherein said nucleotide sequence encodes for a protein having a biological activity of the human IL-11 receptor.
3. The polynucleotide of claim 1 wherein said nucleotide sequence is operably linked to an expression control sequence.
4. The polynucleotide of claim 2 comprising the nucleotide sequence of SEQ ID NO:1 from nucleotide 803 to nucleotide 1999.
5. The polynucleotide of claim 2 comprising the nucleotide sequence of SEQ ID NO:1 from nucleotide 803 to nucleotide 1828 or a fragment thereof.

6. The polynucleotide of claim 2 comprising the nucleotide sequence of SEQ ID NO:1 from nucleotide 1907 to nucleotide 1999 or a fragment thereof.

7. The polynucleotide of claim 1 comprising the nucleotide sequence of SEQ ID NO:1 from nucleotide 734 to nucleotide 1999.

8. The polynucleotide of claim 1 comprising the nucleotide sequence of SEQ ID NO:1 from nucleotide 1067 to nucleotide 1828.

9. The polynucleotide of claim 1 comprising the nucleotide sequence of SEQ ID NO:1 from nucleotide 1067 to nucleotide 1999.

10. A host cell transformed with the polynucleotide of claim 3.

11. The host cell of claim 8, wherein said cell is a mammalian cell.

12. A process for producing a human IL-11R protein, said process comprising:

(a) growing a culture of the host cell of claim 10 in a suitable culture medium; and

(b) purifying the human IL-11R protein from the culture.

13. An isolated human IL-11R protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO:2;
- (b) the amino acid sequence of SEQ ID NO:2 from amino acids 24 to 422;
- (c) the amino acid sequence of SEQ ID NO:2 from amino acids 24 to 365;
- (d) the amino acid sequence of SEQ ID NO:2 from amino acids 391 to 422;
- (e) the amino acid sequence of SEQ ID NO:2 from amino acids 102 to 422;
- (f) the amino acid sequence of SEQ ID NO:2 from amino acids 102 to 365; and
- (g) fragments of (a)-(f) having a biological activity of the human IL-11 receptor.

14. The protein of claim 13 comprising the amino acid sequence of SEQ ID NO:2.

15. The protein of claim 13 comprising the sequence from amino acid 24 to 365 of SEQ ID NO:2.

16. A pharmaceutical composition comprising a protein of claim 13 and a pharmaceutically acceptable carrier.

17. A protein produced according to the process of claim 12.
18. A composition comprising an antibody which specifically reacts with a protein of claim 13.
19. A method of identifying an inhibitor of IL-11 binding to the human IL-11 receptor which comprises:
- (a) combining a protein of claim 13 with IL-11 or a fragment thereof, said combination forming a first binding mixture;
 - (b) measuring the amount of binding between the protein and the IL-11 or fragment in the first binding mixture;
 - (c) combining a compound with the protein and the IL-11 or fragment to form a second binding mixture;
 - (d) measuring the amount of binding in the second binding mixture; and
 - (e) comparing the amount of binding in the first binding mixture with the amount of binding in the second binding mixture;
- wherein the compound is capable of inhibiting IL-11 binding to the human IL-11 receptor when a decrease in the amount of binding of the second binding mixture occurs.
20. The method of claim 19 wherein the first and second binding mixture comprise gp130 or a fragment thereof capable of binding to the protein of claim 13 or the IL-11 or fragment used therein.

21. An inhibitor identified by the method of claim 19.
22. A pharmaceutical composition comprising the inhibitor of claim 21 and a pharmaceutically acceptable carrier.
23. A method of inhibiting binding of IL-11 to the human IL-11 receptor in a mammalian subject, said method comprising administering a therapeutically effective amount of a composition of claim 22.
24. A method of inhibiting binding of IL-11 to the human IL-11 receptor in a mammalian subject, said method comprising administering a therapeutically effective amount of a composition of claim 16.
25. A method of inhibiting binding of IL-11 to the human IL-11 receptor in a mammalian subject, said method comprising administering a therapeutically effective amount of a composition of claim 18.
26. A method of treating or preventing loss of bone mass in a mammalian subject, said method comprising administering a therapeutically effective amount of a composition of claim 22.
27. A method of treating or preventing loss of bone mass in a mammalian subject, said method comprising administering a therapeutically effective amount of a composition of claim 16.

28. A method of treating or preventing loss of bone mass in a mammalian subject, said method comprising administering a therapeutically effective amount of a composition of claim 18.

29. An isolated polynucleotide comprising a nucleotide sequence capable of hybridizing under stringent conditions to polynucleotide of claim 4.

30. An isolated polynucleotide comprising a nucleotide sequence encoding a peptide or protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO:2;
- (b) the amino acid sequence of SEQ ID NO:2 from amino acids 24 to 422;
- (c) the amino acid sequence of SEQ ID NO:2 from amino acids 24 to 365;
- (d) the amino acid sequence of SEQ ID NO:2 from amino acids 391 to 422;
- (e) the amino acid sequence of SEQ ID NO:2 from amino acids 112 to 422;
- (f) the amino acid sequence of SEQ ID NO:2 from amino acids 112 to 365; and
- (g) fragments of (a)-(f) having a biological activity of the human IL-11 receptor.

FIG. 1/1

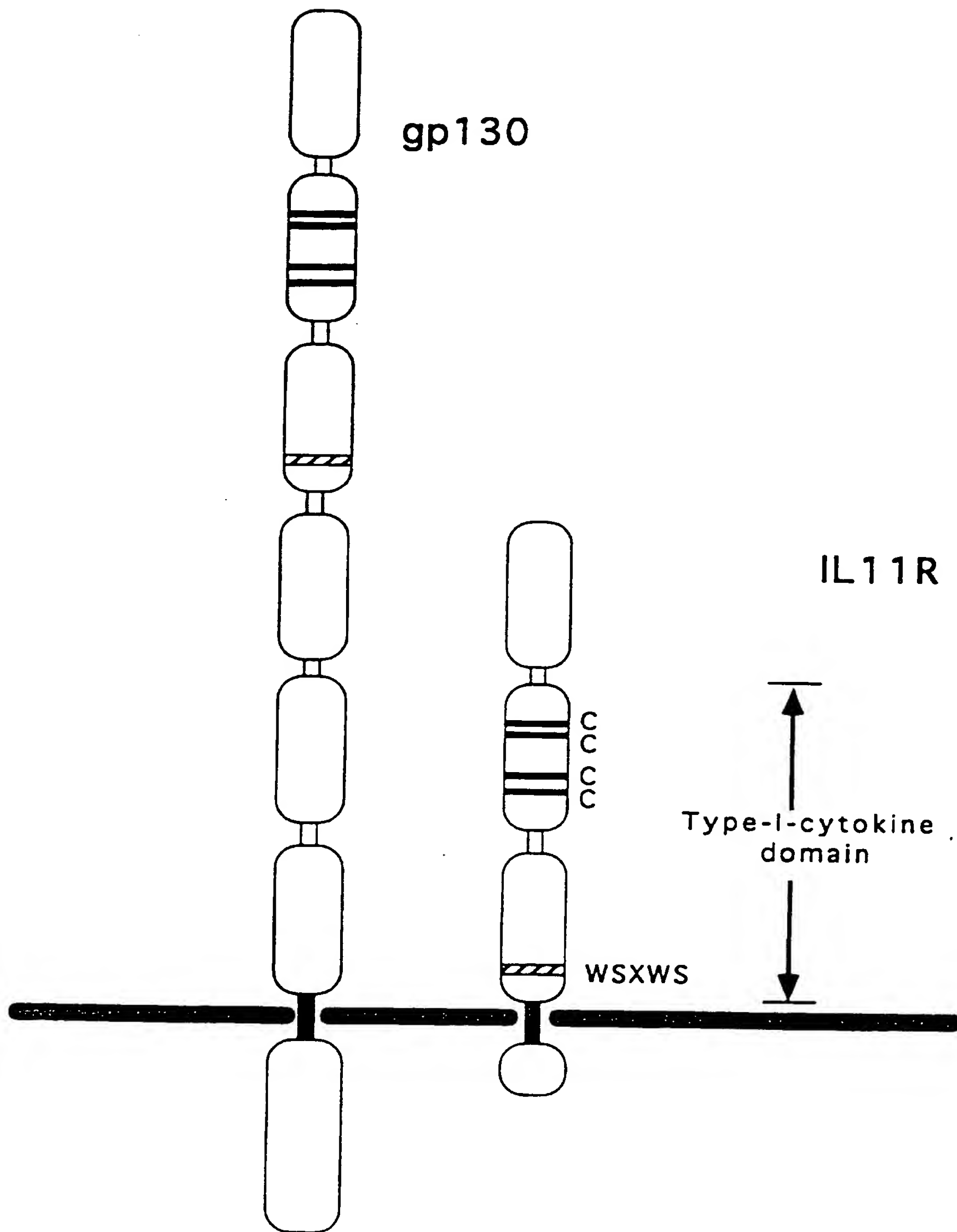
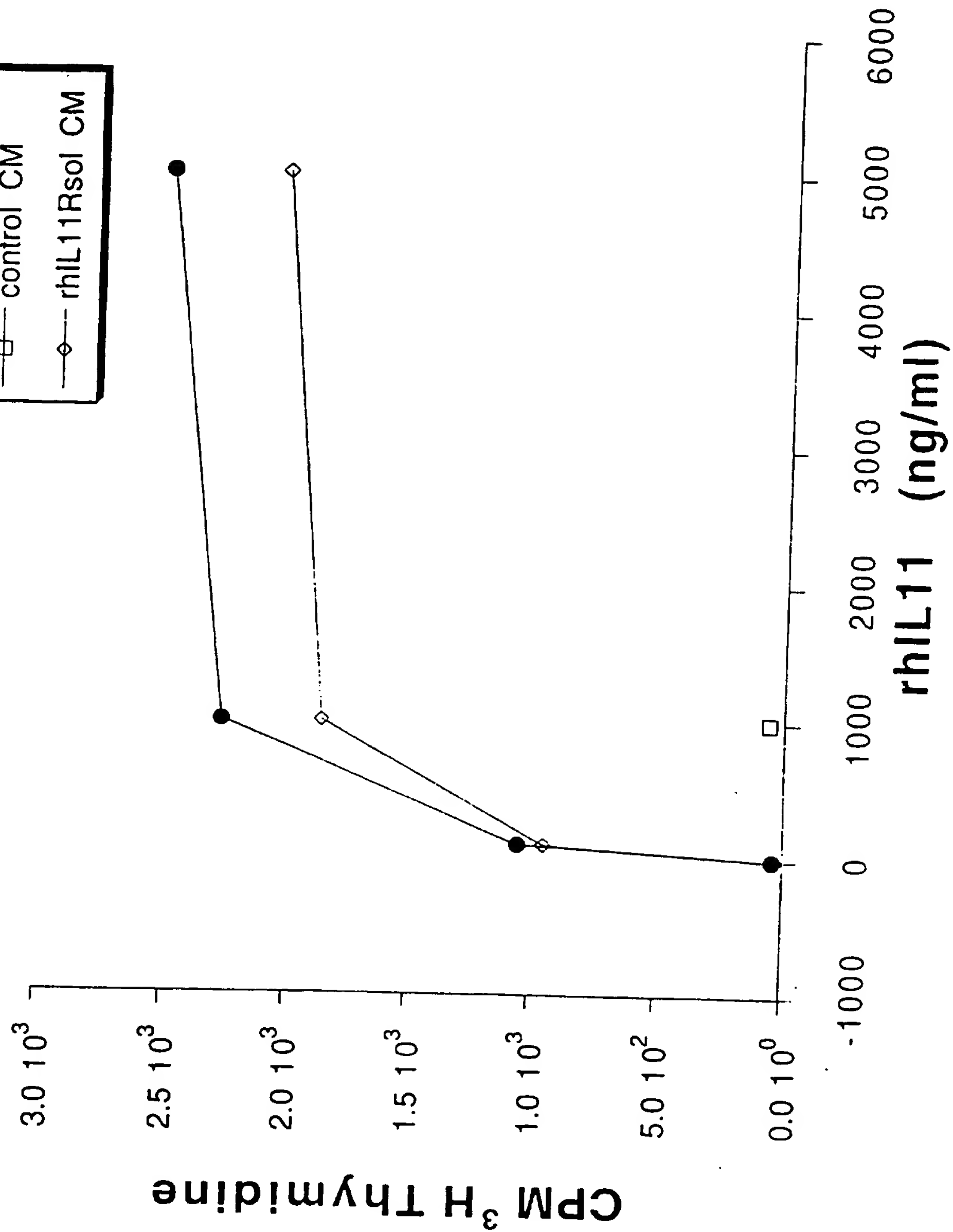


FIGURE 2



INTERNATIONAL SEARCH REPORT

International Application No
PCT/US 95/15400

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 C12N15/12 C07K14/715 A61K38/17 C07K16/28 C12N5/10
A61K39/395 G01N33/68

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C07K C12N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>DEVELOPMENTAL BIOLOGY, vol. 166, 1994, pages 531-542, XP002000295 H. NEUHAUS ET AL: "Et12, a novel putative type-I cytokine receptor expressed during mouse embryogenesis at high levels in skin and cells with skeletogenic potential" *see the whole document especially figure 3 page 535 *</p> <p style="text-align: center;">---</p> <p style="text-align: center;">-/--</p>	1,4-9, 29,30

☒ Further documents are listed in the continuation of box C.

☐ Patent family members are listed in annex.

* Special categories of cited documents :

A document defining the general state of the art which is not considered to be of particular relevance

E earlier document but published on or after the international filing date

L document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

O document referring to an oral disclosure, use, exhibition or other means

P document published prior to the international filing date but later than the priority date claimed

T later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

X document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

Y document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

Z document member of the same patent family

Date of the actual completion of the international search

3 April 1996

Date of mailing of the international search report

23. 04. 96

Name and mailing address of the ISA

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Fax: (+ 31-70) 340-3016

Authorized officer

Le Cornec, N

INTERNATIONAL SEARCH REPORT

International Application No
PCT/US 95/15400

C.(Continuation) D CUMENTS CONSIDERED TO BE RELEVANT		
Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>EMBO JOURNAL, vol. 13, no. 20, 17 October 1994, EYNSHAM, OXFORD GB, pages 4765-4775, XP002000296 D.J. HILTON ET AL: "Cloning of a murine IL-11 receptor alpha-chain ; requirement for gp130 for high affinity binding and signal transduction" see page 4766, right-hand column, line 9 - line 37 see page 4769, right-hand column, line 17 - line 20 see figure 1</p> <p style="text-align: center;">---</p>	1,4-9, 29,30
P,X	<p>BLOOD, vol. 86, no. 7, 1 October 1995, pages 2534-2540, XP002000297 M. CHEREL ET AL: "Molecular cloning of two isoforms of a receptor for the human hematopoietic cytokine Interleukin-11" see the whole document</p> <p style="text-align: center;">---</p>	1,2,4-9, 29,30
A	<p>EUROPEAN JOURNAL OF IMMUNOLOGY , vol. 24, no. 1, January 1994, pages 277-280, XP002000298 M. FOURCIN ET AL: "Involvement of gp130/interleukin-6 receptor transducing component in Interleukin-11 receptor" see page 278, right-hand column - page 279</p> <p style="text-align: center;">---</p>	19,20
A	<p>BIOFACTORS, vol. 4, no. 1, December 1992, pages 15-21, XP002000299 YU-CHUNG YANG ET AL: "Interleukin-11 and its receptor" see page 17 - page 18</p> <p style="text-align: center;">-----</p>	13-15,19